Importance and Utilization of the Genetic Resources of Cultivated Species

Universidad Nacional Agraria La Molina 50th Anniversary of the Escuela de Post Grado "Research for a Sustainable Development" September 16-17, 2008

Candice Gardner, USDA-ARS, Ames, IA





Food security depends upon germplasm collections.

http://www.fao.org

In agriculture, the most important resources are soil, water, air, and germplasm collections.

Plant genetic resources (PGR) have been utilized over the millennia to improve the human condition. Development of crops that provided stable supplies of food, feed, fiber and fuel offered alternatives to nomadic existence, and enabled societies to develop and flourish around the world.

The development of improved crops and cropping systems, and the increased availability of materials essential for sustenance, growth and development made possible the rapid evolution of human intellectual capacity, discovery and invention. As a result, the nature of human society and human behavior changed.

Conservation and utilization of PGR have made possible the continuous varietal improvement necessary to provide solutions for agricultural production challenges, the development of new crops and new uses, and improvements in health and nutrition.

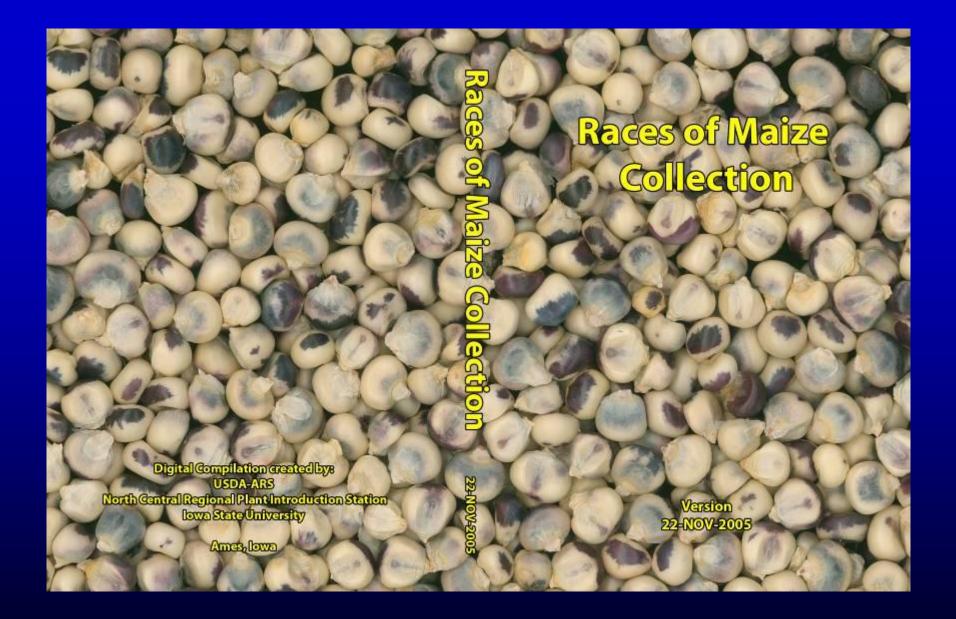
Growing global energy needs coupled with the increasing demands for food and other plant-based resources have highlighted the critical importance of PGR now more than ever. Demand for well-documented PGR increases annually.

The Four F's

- Food
- Feed
- Fiber
- Fuel



The World's #1 Crop





The Other World's #1 Crop



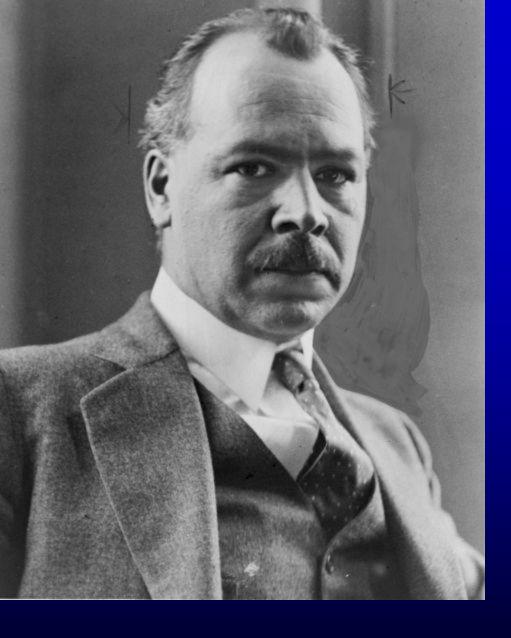




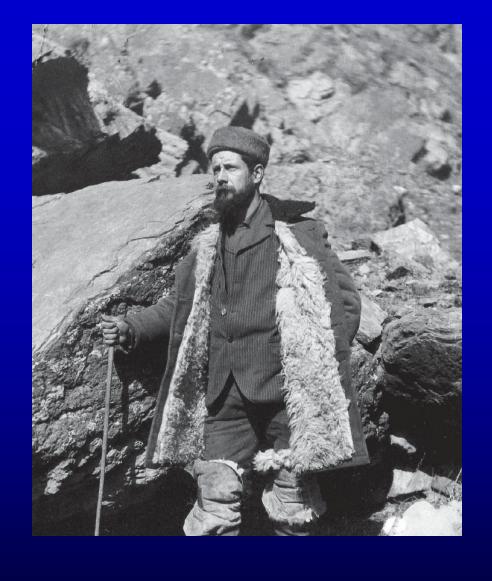




The Early Plant Explorers



Nikolai I. Vavilov is recognized as the foremost plant geographer of contemporary times. To explore the major agricultural centers in this country and abroad, Vavilov organized and took part in over 100 collecting missions, including those to Iran (1916), the United States, Central and South America (1921, 1930, 1932), the Mediterranean and Ethiopia (1926-1927). He and his colleagues were the first to thoroughly collect potato germplasm in the Andes.



1910, the Office of Foreign **Seed and Plant Introduction** of the United States sent Frank N. Meyer, one of world's most outstanding collectors, to the Sinkiang **Province of China. He was** also to go to Europe, Russia, and Tibet. During 1910 to 1911, Meyer explored Irkutsk, Kashgar, Yarkand, and from there to Ak-Suand and on to Kul'dzhe, always in search of hardy crop plants and their wild relatives that could tolerate abiotic stresses.

Potato Germplasm Collection Development

The dedication and effort of pioneer taxonomist Carlos Ochoa formed the foundation of the CIP-held collection of potato genetic resources. This Peruvian scholar, whose crusade to find wild species in the Andes began more than 40 years ago, has discovered 80 different species—about one-third of all the wild potatoes known to exist.





© II-K Forian Centanh

Drs. Salas & Spooner Collecting Wild Germplasm



"THE DISAPPEARANCE OF OLD VARIETIES, the

landraces of crop plants, and their wild progenitors could eventually be recognized as the great sleeper issue in the last decades of the 20th century.

It is difficult for us to visualize a scenario more profound in its implications, yet less appreciated by funding institutions, governments, and the general public, than that entailed in the mass elimination of a large number of plant species that has taken place and continues to take place in the centers of their diversity."

(Qualset and Shands, 2005).

Recent Threats to Global Food Production

- Insect and Disease Pests
 - Wheat Stem Rust Reemerges
- Fertility Costs and Availability
- Energy Costs
- Water
- Stable Supply of Adapted Cultivars
- Land Use / Land Loss

Plant Germplasm Collection Development Today

- Collection expeditions
- Germplasm originators

- Exchange between institutions
 - Within National Institutions
 - Between Nations, governed by the ITPGRFA

Successful Conservation & Utilization of PGR Depends On:

- Targeted acquisition
- Documentation
- Effective maintenance and regeneration programs
- Thorough evaluation and characterization
- Effective use of associated information
- Distribution of PGR to research communities

Last but not least...

• Pre-breeding activities to facilitate utilization.

 Thorough understanding of the phenotypic and genetic variability of a crop and its wild relatives, their adaptation, life forms, breeding systems, traits and biological properties supports PGR conservation activities, and is essential to realize their potential for contribution.

Current Status: Plant Genetic Resources

- An estimated 6 million samples in genebanks world-wide; of these, 1.5 million are probably unique.
- About 10% of the samples are held by the IARCs.
- Most samples are in genebanks controlled by national governments.

- Large national genebanks in the U.S., Canada, Australia, Japan, S. Korea, India, China, Brazil, Russia, Germany, South Africa.
- Public gardens, NGOs, universities, and companies also hold thousands of samples.

The Global Crop Diversity Trust is an independent international organization which exists to help ensure the conservation and availability of crop diversity for food security worldwide. It was established through a partnership between the United Nations Food and Agriculture Organization (FAO) and the Consultative Group on **International Agricultural Research** (CGIAR).

Annex 1 Crops of the ITPGRFA

The crops listed include: breadfruit, asparagus, oat, beet, brassicas (the cabbage family including broccoli and cauliflower), pigeon pea, chickpea, citrus, coconut, aroids (including taro and cocoyam), carrot, yams, finger millet, strawberry, sunflower, barley, sweet potato, grass pea, lentil, apple, cassava, banana/plantain, rice, pearl millet, beans, pea, rye, potato, eggplant, sorghum, triticale, wheat, faba bean, cowpea, maize and more than 80 forage species from 30 different genera.

The U.S. National Plant Germplasm System (NPGS)

- "Base collection"; preservation research.
- GRIN database: www.ars-grin.gov
- Acquisition via plant exploration and exchange.
- Germplasm quarantine activities now part of APHIS; ARS conducts related research.

- 26 active US sites manage clonally and seed-propagated collections.
- Conduct associated research.
- Crop Germplasm
 Committees;
 university, NGO,
 industry cooperators,
 and ARS.

National Plant Germplasm System Types of germplasm

- Wild crop relatives
- Landrace collections
- Genetic stocks
- Domestic breeding lines
- Heirloom accessions
- Cultivars
- Plant mycosymbionts

US Regional Plant Introduction Station Functions

Conserve Plant Genetic Diversity.

- Encourage Use of Germplasm.
- Conduct Research to Improve Genetic Resource Management Programs.
- Generate Information to Better Target Germplasm Use by the User Community.















Activities of the NPGS Integrate Data Acquisition to Enhance their Efficiency and Collection Value

Acquisition
Documentation
Maintenance & Regeneration
Characterization
Evaluation
Distribution

Germplasm collection now utilizes GIS technology

Participants in a 2001Kazakhstan expedition:

Richard Hannan
Stephanie L. Greene
Alexandr Afonin
Nickolai Dzubenko

Institutes involved

- USDA, ARS National Plant Germplasm System
- Kazakhstan Institute of Agricultural Science-Aral Sea Experiment Station
- N.I. Vavilov Institute of Plant Industry, Russia
- University of St. Petersburg, Russia

Results

- Trip covered 1180 km (~ 730 miles)
- Collections were made at 89 sites
- 432 accessions were collected:
 - 138 Medicago
 - 86 Legumes (Astragalus, Vicia, Lathyrus, Trifolium)
 - 48 Grasses
 - 58 Wild onions (2-3 species)
 - 102 Other horticulture (wild forms of hops, medicinal species, lettuce, ornamentals, cultivated garlic, apples)

Hypericum Collection in the Ozarks



Wild *Helianthus* Collection in California



US NPGS System Holdings

- 26 Active Sites + the NCGRP (Ft. Collins)
 + the NGRL
- 33 Designated Collections
- 508,622 Accessions as of September 13, 2008
- 20-25% of Accessions Distributed Yearly
- 2,126 Genera; range of 1-583 per site
- 13,111 Species; range of 1-3,058 per site
- 235 Families Represented

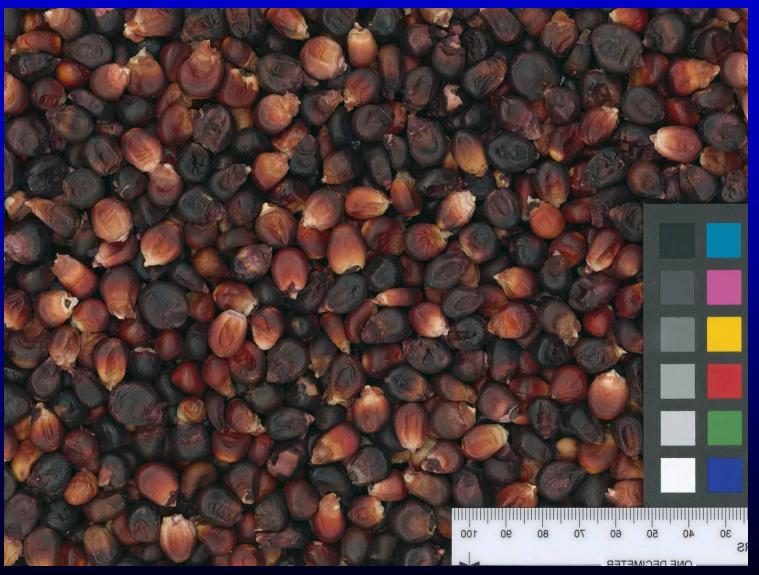
Activities of the NPGS Integrate Data Acquisition to Enhance their Efficiency and Collection Value

Acquisition
Documentation
Maintenance & Regeneration
Characterization
Evaluation
Distribution

Crop Evolution Lab Records

	4		
			dewest 4348 4.5 miles south of Ahuacatlan and 2.7 mi south of Jala turnoff along Hex. 15 in Wayarit State
	4736	Panicum hirticoule	dewest 4348 4.3 miles sown of the actual and a string south of season as the
	4737	Panieum sonorum	dewed 4362 1 miles 30 with east of Circlad Obregon along HEX. 15 in Sonora State
	4738	Panicum sonorum	dewed 4365 // miles east of Navajoa along & Alamos road Sonora, Hex
	4739	Panicum sonorum	deWest 4366
	4740	Panicum Sonorum	dewed 4367 "
	4741	Panicum Sonorum	dewed 4373 20.4 miles N.W. of Alamos and 136 miles E of Navojoo. Sonora, Hex. along the Alamos Road
	4742	Panicum songrum	dewed 4374 23 miles NW of Alamos and 11 mile East of Navojoa, Sonora, Hex along Alamos Rd. on easternside of Cerro Prieto
	4743	Paricum sonorum	dewed 4375 23 miles N.W. of Alamos, Sonora, Hex. along road to Alamos past base of Cerro Prieto.
	4744	Panicum sonorum	de Wet 4400 among Ruinos de Ixtlan. 21 mi. JE of little forwar of Ixtlan del Rio Nayarit.
	4745	Pariam sonoum	de Wed 4401 2.7 mi. SE of Ixtlas du Rio, Nayarit just off Mex. 15.
	4746	Panieum sonorum	dewed 4402 115 mi. east of Ixtlan, Nayarit in Jalisco along Hex. 15.
RECO	4748	Paricum sonorum	dewas 4405 15.4 mi. 5E of Tequila and 1.7 from El Arenal, Jalisco along Hes. 15.
	4749	Panicum Straminaum	dewed 4753 3.3 mi. south of Casis de los Chinas of 57.8 south of Santa Ona in Sonora State, Hex.
	4750	Paneum straniveum	dewed 4395 B.2 mi. SE of Escuinapa and 66.8 of Mazatlan, Sinaloa, along road, Alex. 15
	4751	Panicum stramineum	ou wet 4348 78.3 mi SE of Hazattán ard 24.7 mi SE of Escuinapa, along Hex. 15 to Tepic
	4752	Panicum adopersum	wet 4415 8mi, N. of US 77 + US 83 junction in Harlington, Texas

Acquisition of Images





Genebank Information Databases

A major function of a genebank is to manage the information associated with collections and provide them in a usable format, both for collection management and to facilitate utilization.

Examples of germplasm information databases would include GRIN, SINGER, and EURISCO, among others.

- Information associated with the collections increases their value to researchers and to genebank managers, enables better targeting to meet research objectives.
- Automation of data collection and transfer activities improve genebank resource use efficiency.
- IT and IM Tech transfer between sites and between researchers is increasingly critical for successful germplasm conservation and utilization.
- Interoperability with genomic databases is a high priority
- Stakeholder input is key to the future utility of PGR information delivery systems.

Use of Existing Information to Assist in Collection Development Strategies

too much information can be overwhelming...

not enough

Readily accessible, useful information is enabling

GRIN-Global

GRIN-Global is a project whose mission is to create a new scalable, version of the GRIN system suitable for use by any interested genebank in the world. It is being developed in a joint effort with the Global Crop Diversity Trust, Bioversity International, and the Agricultural Research Service of the USDA. Replacement of the current GRIN system for NPGS use with the GRIN-Global system is scheduled for the last quarter of 2010.

http://cool.ars-grin.gov/wiki/index.php/Main_Page

http://cool.ars-grin.gov/gringlobal/forums/

Project Goals

 To provide the world's crop genebanks with a powerful, flexible, easy-to-use global plant genetic resource (PGR) information management system that will constitute the keystone for a sustainable, rational, efficient, and effective global network of genebanks to permanently safeguard PGR vital to global food security, and to encourage the use of PGR by researchers, breeders, and farmer-producers.

 The database and interface(s) will be designed to accommodate both commercial and open-source programming tools, to be database-flexible, and to require no licensing fees for genebank use. This will enable institutions with limited IT resources, as well as better-supported genebanks, to adopt GRIN-Global. The database will be deployable on local standalone computers at sites with limited computational capabilities, as well as at networked sites.

Genetic resources flow chart

Accession in genebank

Morphological data

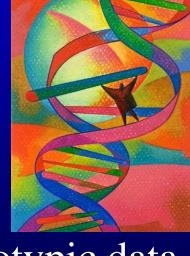
Phenotypic data

Disease

Nutrition

Yield components





Genotypic data

Accession Utilization

Activities of the NPGS Integrate Data Acquisition to Enhance their Efficiency and Collection Value

Acquisition
Documentation
Maintenance & Regeneration
Characterization
Evaluation
Distribution

The objective of Maintenance and Regeneration is to provide high quality seed and plant materials which are true to the original genetic profile.

Viability & Preservation Research

- The main research efforts in this area within the USDA are conducted at the NCGRP in Ft. Collins, CO, but also at several other sites.
- Lipid chemistry is known to impact effective storage practices.
- Cryopreservation techniques are being successfully developed for woody species; apple, willow, and ash are recent examples.
- Tissue culture is essential to both micropropagation of species such as strawberry or potato; it also provides for resolution of some phytosanitary issues.

- •Controlled pollination is necessary for cross-pollinated crops, and is accomplished by hand- or insect-mediated methods.
- •Clonal and micropropagation methods require controlled conditions for the production of pathogen-free propagules
- •ALL CONTAMINATION IS BAD, REGARDLESS OF THE SOURCE





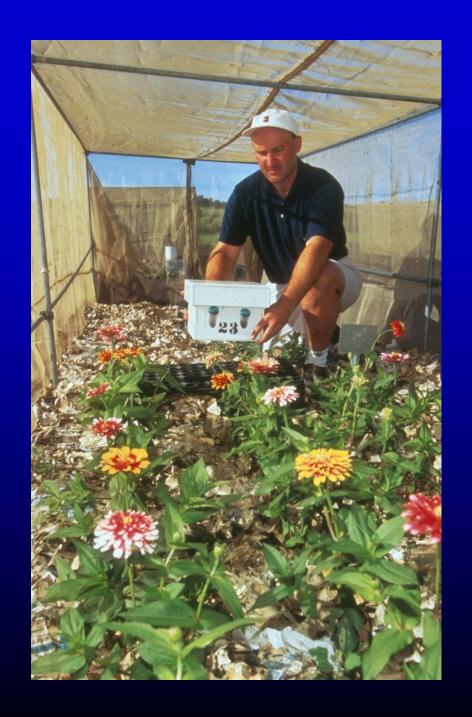




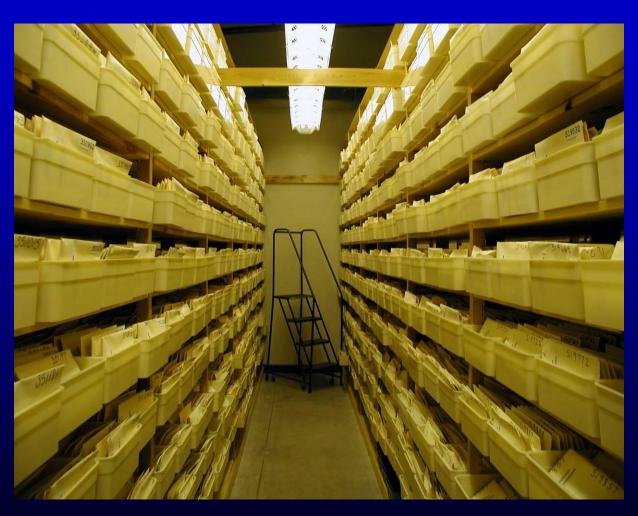








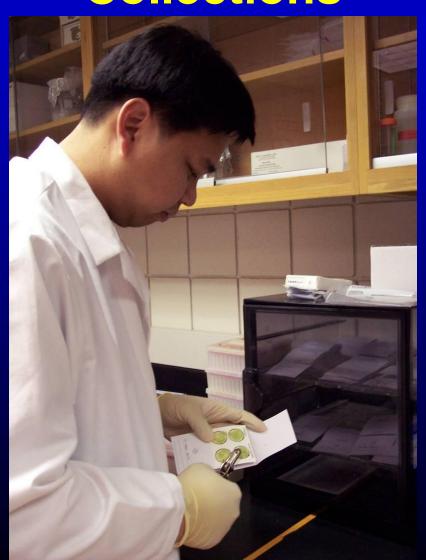
Seed Storage at the National Small Grains Collection, Aberdeen, ID





Activities of the NPGS Integrate Data Acquisition to Enhance their Efficiency and Collection Value Acquisition Maintenance & Regeneration Characterization **Evaluation Documentation Distribution**

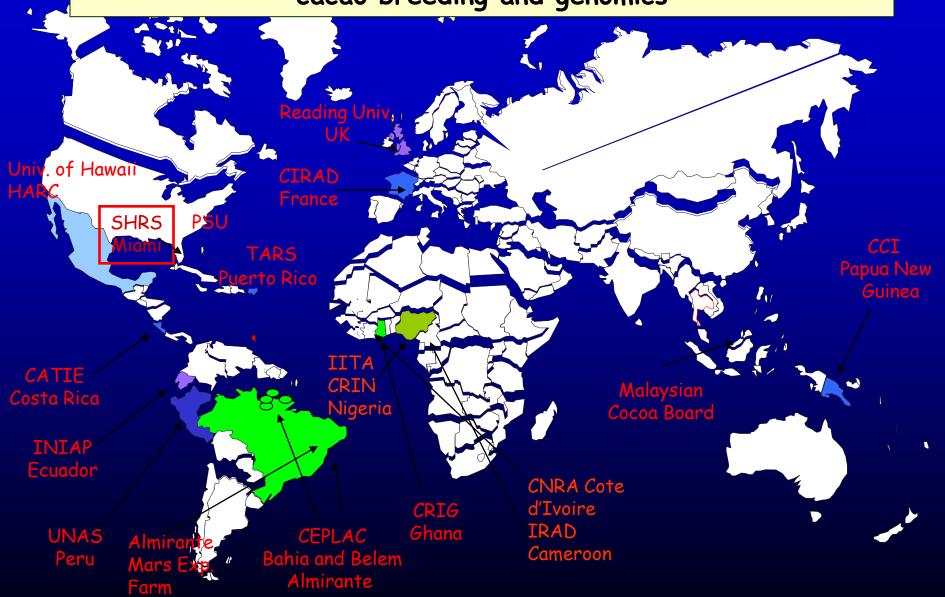
Use of DNA Capture Cards for Molecular Characterization of Collections



Sequencing, annotation, and database management of the genome of Theobroma cacao

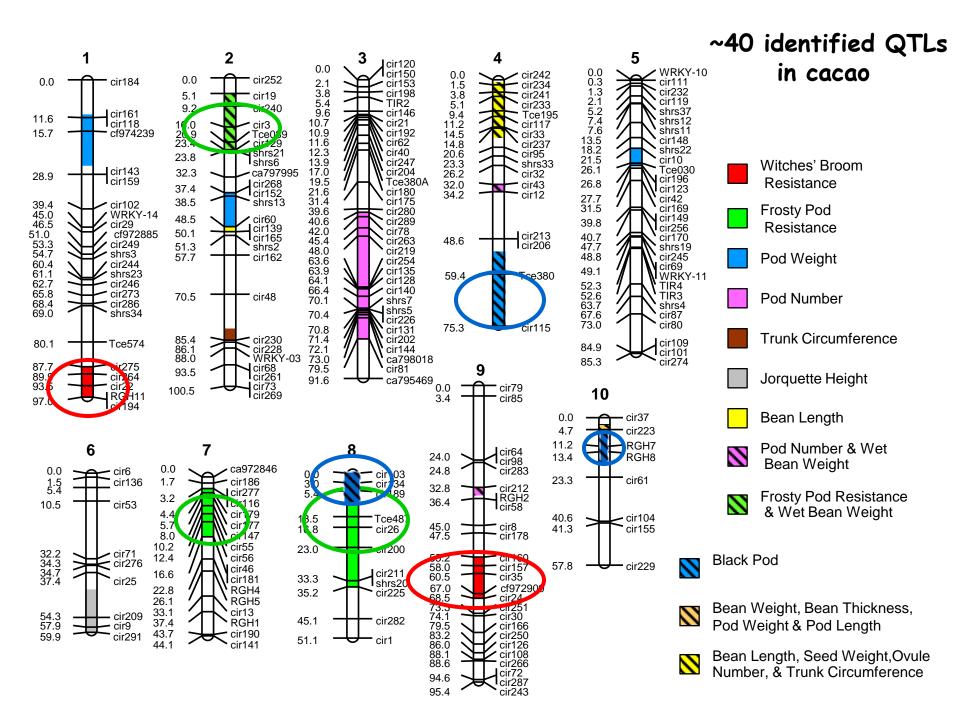
- A composite linkage map from the combination of three crosses made from commercial clones of cacao, *T. cacao* L. Tropical Tree Genomics and Genetics. DOI 10.1007/s/2042-008-9011-4. Brown J. Steven Brown, Robert T. Sautter, Cecile Olano, James W. Borrone, David N. Kuhn, and Raymond J. Schnell. 2008.
- Development of a Marker Assisted Selection Program for Cacao. Phytopathology Vol 97, No 12, 1665-1669. Schnell, R. J., D. N. Kuhn, J. S. Brown, C. T. Olano, and J. C. Motamayor. 2007.

SCA and other formal and informal collaborative projects for cacao breeding and genomics



Current field evaluations

Research Stations	Genotypes	
South and Central America		
CATIE, Costa Rica	4,000	
INIAP, Ecuador	12,500	
MARS Farm Almirante, Brazil	4,500	
IBE, Peru	500	
West Africa		
CNRA, Cote d'Ivoire	1,500	
CRIG, Ghana	5,000	
CRIN, Nigeria	500	
IRAD, Cameroon	300	
Asia		
CCI, PNG	2,000	
USA		
Miami (quarantine and nursery)	250	
Mayaguez (germplasm collection)	350	
Hawaii (selection on Oahu and Hawaii)	1,600	
Total genotypes for project	33,000	



Number of plants with the favorable alleles for disease resistance at the two QTL for WB and three QTL for FP in families currently under evaluation at CRIG in Ghana.

n	WB	BP	WB+BP
79	19	6	2
90	28	19	6
109	34	11	6
119	11	1	0
15	4	2	1
412	96	39	15
	79 90 109 119	79 19 90 28 109 34 119 11 15 4	79 19 6 90 28 19 109 34 11 119 11 1 15 4 2

Where are we going next?

The development of SNP markers in genes involved widisease resistance, productivity, and quality.

To do this we need to sequence the genome and produce the genetic linkage map.

The MAS breeding program would be greatly enhance accelerating the delivery of improved, productive, dise resistant cultivars to cocoa farmers.

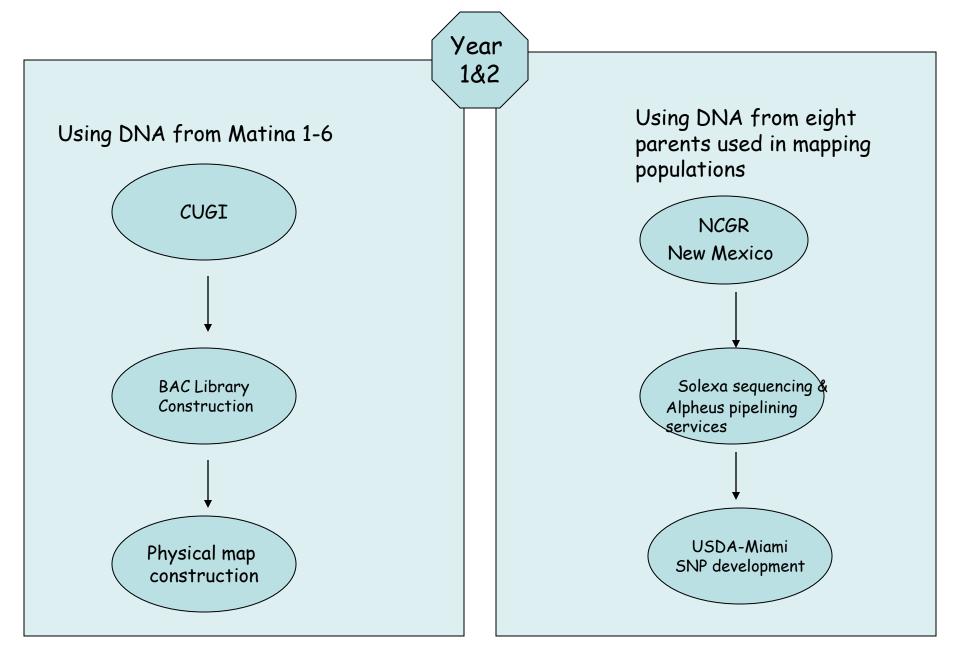
Choice field location on Kona side of Hawaii



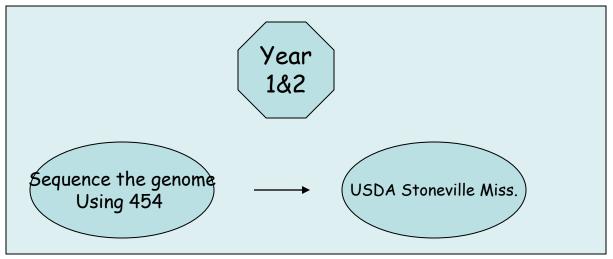
Sequencing the *T. cacao* genome

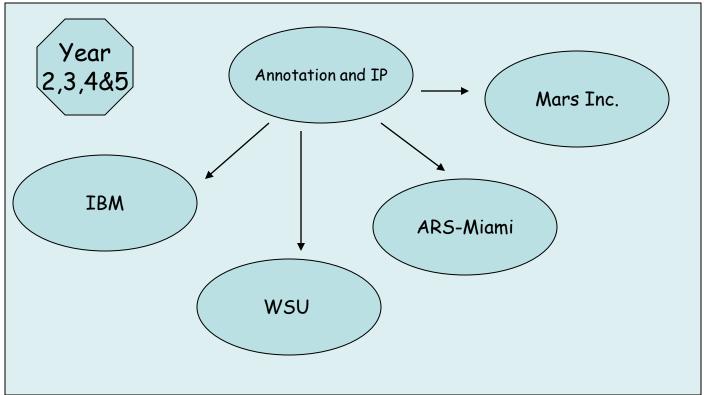
Species	Genome si	ze
Yeast	12 Mb	
Arabidopsis	119 Mb	
Theobroma cacao	415 Mb	
Rice	430 Mb	
Dog	2,400 Mb	
Human	3,300 Mb	

Whole genome sequencing workflow



Whole genome sequencing workflow





Activities of the NPGS Integrate Data Acquisition to Enhance their Efficiency and Collection Value

Acquisition
Documentation
Maintenance & Regeneration
Characterization
Evaluation
Distribution

Adaptation Criteria

Ames in the Wintertime



Ames, IA in the Summertime





50%
Highland
Tropical
Maize –

Unadapted to the MidWest



GEM Project
Has
Successfully
Introgressed

Exotic Germplasm –

190 Lines
Released with
Adaption to
the Midwest or
Southeast

Activities of the NPGS Integrate Data Acquisition to Enhance their Efficiency and Collection Value

Acquisition
Documentation
Maintenance & Regeneration
Characterization
Evaluation
Distribution

CRIS Project Search for US - NCR

A search of the Cooperative Research Information System (CSREES and ARS Projects) resulted in identification of 528 projects using plant genetic resources on September 4, 2008.

1,459,816 samples were distributed by the NPGS from 2000-2006, or an average of 25% of collection holdings annually. Between 30 and 40% were distributed to international recipients.

Next three slides courtesy of Luciano Nass, EMBRAPA-Labex USA

BRAZIL – SUGARCANE in 2008

(1,000 tons)

	SUGAR AND ETHANOL INDUSTRY			
TOTAL	TOTAL	SUGAR	ETHANOL	OTHERS
710,280	558,720	240,890	317,830	151,560
		43%	57%	

Source: CONAB, 2008

BREEDING AND GENETIC ENGINEERING

- Sugarcane varieties = interspecific hybrids
 - Introduced to Brazil 14th century
 - Complex genetically (2n = 70-120) with large DNA content.



Saccharum officinarum (2n = 80)



Saccharum spontaneum (2n = 40-128)



Saccharum robustum (2n = 60-205)



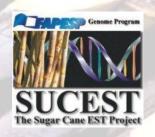
Saccharum barberi (2n = 111-120)

BREEDING AND GENETIC ENGINEERING

- Historically Brazil has had an intensive sugarcane breeding program
 - 550 varieties developed to date
 - 51 varieties released since 1995
 - 20 varieties account for 70% of total planted area
- Sugarcane Genome Project

(http://sucest.cbmeg.unicamp.br/en)

- Understand commercial cultivar origin
- Identification of diversity and genetic variability
- Introgression and QTLs identification (Quantitative Trait Loci)
- Diagnostics for disease resistance / tolerance
- Structural and functional genomics





Number of published papers on sugarcane



Number of published papers on ethanol extracted from biomass



Source: Fapesp (2008)

Early Guayule Production for Natural Rubber











Courtesy of Terry Coffelt, USDA-ARS, Maricopa, AZ

Guayule – A Commercial Crop for Production of High quality, Natural Rubber Latex for Hypoallergenic Products



The Zea nicaraguensis Saga















Education for a Sustainable Future

